

SEQUENCE LISTING

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PABO, CARL

<120> CONTEXT SENSITIVE PARALLEL OPTIMIZATION OF ZINC FINGER
DNA BINDING DOMAINS

<130> 51588.62031 US

<140> 10/532,258
<141> 2005-04-21

<150> PCT/US03/034010
<151> 2003-10-23

<150> 60/420,485
<151> 2002-10-23

<150> 60/466,889
<151> 2003-04-30

<160> 53

<170> PatentIn Ver. 3.3

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peptide

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<223> Variable amino acid and this region may encompass
2 to 4 residues

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<221> MOD_RES
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<223> Variable amino acid

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<221> MOD_RES
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<223> Variable amino acid and this region may encompass
3 to 5 residues

<400> 1

Cys Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1 5 10 15

Xaa Xaa His Xaa Xaa Xaa Xaa Xaa His

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Asp Arg Ser Ser Leu Thr Arg
1 5

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Gln Gly Gly Asn Leu Val Arg
1 5

<210> 4
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<400> 4
Gln Ala Ala Thr Leu Gln Arg
1 5

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<400> 5
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gaa

3

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<400> 7
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Asp Arg Ser Ser Thr Arg Gln Gly Gly Asn Val Arg Gln Ala Ala Thr
1 5 10 15

Gln Arg

<210> 9
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9

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<400> 10
Arg Lys Asp Ser Val Arg Gln Ser Gly Asp Arg Arg Asp Cys Arg Asp
1 5 10 15

Ala Arg

<210> 11
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Ala Ser Ala Asp Thr Arg Asn Arg Ser Asp Ser Arg Thr Ser Ser Asn
1 5 10 15

Lys Lys

<210> 13
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oligonucleotide

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25

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oligonucleotide

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25

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oligonucleotide

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25

<210> 17
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oligonucleotide

<400> 17
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Gln Arg

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Gln Arg

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<400> 20
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Gln Arg

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<400> 21
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Gln Arg

<210> 22
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Thr Arg

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Gln Arg

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Asp Ser Pro Thr Arg Arg Gln Arg Thr Asn Ile Arg Gln Arg Asn Thr
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Gln Arg

<210> 25
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<400> 25
Asp Arg Pro Thr Arg Arg Gln Gly Ala Asn Arg Arg Gln Ala Asn Thr
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Gln Arg

<210> 26
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<400> 26
Asp Ser Pro Thr Arg Arg Gln Asn Thr Asn Asn Arg Gln Ala Asn Thr
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Gln Arg

<210> 27
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<400> 27
Asp Ser Pro Thr Arg Arg Gln Lys Pro Asn Asp Arg Gln Gly Asn Ser
1 5 10 15

Ile Arg

<210> 28
<211> 18
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<400> 28

Asp Ser Pro Thr Arg Arg Gln Ser Thr Asn Asn Arg Gln Gly Asn Ser
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Asn Arg

<210> 29

<211> 18

<212> PRT

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<223> Description of Artificial Sequence: Synthetic peptide

<400> 29

Asp Ser Pro Thr Arg Arg Gln Asn Val Asn Thr Arg Gln Arg Asn Thr
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Gln Arg

<210> 30

<211> 18

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<223> Description of Artificial Sequence: Synthetic peptide

<400> 30

Arg Gln Asp Ile Val Lys Gln Ser Ser Thr Thr Arg Glu Lys Gln Gly
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His Arg

<210> 31

<211> 18

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<400> 31
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Lys Arg

<210> 32
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Lys Arg

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<400> 33
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Lys Arg

<210> 34
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Arg Arg

<210> 35
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Gly Arg

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Lys Arg

<210> 37
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Arg Arg

<210> 38
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Lys Arg

<210> 39

<211> 18

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<400> 39

Arg Gln Asp Ile Val Lys Gln Ser Ser Thr Thr Arg Glu Lys Gln Gly
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His Arg

<210> 40

<211> 18

<212> PRT

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<400> 40

Arg Ser Asp Leu Thr Lys Gln Ser Ser Thr Thr Arg Glu Arg Gln Gly
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Lys Arg

<210> 41

<211> 18

<212> PRT

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<400> 41

Arg Pro Asp Val Gln Lys Gln Ser Gly Thr Val Arg Asp Gln Ser Asn
1 5 10 15

Arg Arg

<210> 42
<211> 18
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<400> 42
Leu Arg Ala Asp Asp Asn Leu Ser Gln Thr Lys Arg Ile Arg Gly Asn
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Val Arg

<210> 43
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<400> 43
Ala Lys Ala Asp Asp Arg Leu Ser Gln Thr Lys Arg Val Lys Ser Asn
1 5 10 15

Arg Arg

<210> 44
<211> 18
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1 5 10 15

Arg Arg

<210> 45
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<400> 45
Leu Arg Ala Asp Asp Arg Leu Ser Gln Thr Lys Arg Val Lys Ser Asn
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Arg Arg

<210> 46
<211> 18
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<400> 46
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1 5 10 15

Ala Arg

<210> 47
<211> 18
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<220>
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<400> 47
Leu Arg Thr Asp Asp Arg Leu Ser Gln Thr Arg Arg Leu Arg Ser Asn
1 5 10 15

Gly Arg

<210> 48
<211> 18
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peptide

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1 5 10 15

Met Arg

<210> 49
<211> 18
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peptide

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1 5 10 15

Leu Arg

<210> 50
<211> 18
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peptide

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1 5 10 15

Thr Arg

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peptide

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5

10

15

Lys Arg

<210> 52

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Leu Arg

<210> 53

<211> 18

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Ser Arg